



FIGURE 2

```
Motif 1
                                  (1) ----MSWLQYPEGIDPADVQLINNEAFDNGDSEDLSBIKR
                        CDS0689
                                  CDS0689 At
      CDS0689 Medicago trunculata
                     CDS0689 Os
              CDS0689 Ta variant
                     CDS0689 So
           CDS0689 Zm variantrev
              CDS0689 So variant
                 CDS0689 Bn est1
 CDS0689 Eschscholzia californica
                CDS0689 Ga est1
                                  (1) ---MSSILQYPÖVVDAPEVQIMNTAFTNGESEGSLNIKS
(1) MNEVLHIQEARTDSSTDHQIMNTAFTSGESEDSPVVID
(1) ---MSYLQYPÖTLNGQETQINNIAFTNGESEDSTAKKG
                     CDS0689 Pt
        CDS0689 Plumbao zeylanica
         CDS0689 Citrus sinensis
                      Consensus
                                  (1)
                                               E IPEVQIWNNAAFD GDS S AI
                                                            Motif 2
                        CDS0689
                                 (37) -- SWSPLKPLSVRPSDSFESDLSSKENOTPLFENSSVNLS
                                 CDS0689 At
      CDS0689 Medicago trunculata
                                 (32) ---SP-----AVAAVRKGOKEKHRIEVVD-----
                     CDS0689 Os
                                 (32) ------TPVRRGEKENRRAETN-----
(32) ------VPASIRREAEGDKENHREDPDP-----
              CDS0689 Ta variant
                     CDS0689 So
                                 CDS0689 Zm variantrev
             CDS0689 So variant
CDS0689 Bn est1
 CDS0689 Eschscholzia californica
                CDS0689 Ga est1
                    CDS0689 Pt
                                     --F----SAPNLSOSILSOSSIE BLSPSLAEMPHPAK
       CDS0689 Plumbao zeylanica
         CDS0689 Citrus sinensis
                                     SWANLKS-----VYMNGSLESDCSKENLSERLNKSPTSSL
                      Consensus
                                 (41)
                                                    S ESDG KEN P
                                     81
                       CDS 0689
                                 (75)
                                     SPLPIKPLNPNGALENSRLKPNKPNSKQSLDEMAARKSGK
                                     VSITEAPS-----SNK---IVPS-----SNK-
                    CDS0689 At
                                 (69)
     CDS0689 Medicago trunculata
                                 (49)
                                     ---VAAG-----
                    CDS0689 Os
                                 (53)
             CDS0689 Ta variant
                                     _____
                                 (50)
                                 (54) -----
                    CDS0689 So
          CDS0689 Zm variantrev
                                 (55) -----
             CDS0689 So variant
                                (54)
               CDS0689 Bn est1
                                 (67) VSFSTDDP-----EKPH
CDS0689 Eschscholzia californica
                                (75) SPLPIKPLHPNG-----
               CDS0689 Ga est1
                                (72) STASVVKP-----LSSKNVTGNTREPFSAKMKSGVCKE
                   CDS0689 Pt
                                (66) NSSKPAKPLO-----
      CDS0689 Plumbao zeylanica
                                (73)
                                    SPMQK-----
        CDS0689 Citrus sinensis
                                     KSCVPNKPLQVN-----SSVKNSQMKQLK--SVSK
                                (72)
                     Consensus
                                (81)
                                                Motif 3: coiled coil
                               (115) GNDFRDEKKEDEEIEEIQMEISRLSSRLEALREEKAEKTV
                       CDS0689
                    CDS0689 At
                                (94) -----RDEDAE LEEVEKE I GRLSTELES LREEKAEQTA
                                (56) -----RTEDDEIAEIESEIKRLESKLELLRVEKAERKI
    CDS0689 Medicago trunculata
                    CDS0689 0s
                                (57) -----YDŸĒAFIGHIEAEILRLŠSRLHHLRŸSKQPEPN
            CDS0689 Ta variant
                                (50) -----DADAEIARIEAEILRLSSRLHHLRYSKGHDAK
                                (54) -----DYEAEIGHIEAEILRLXSRLHHLRTSKQSEPS
                    CDS0689 So
                                (55) -----ÉEEMRHIEAEILRLSLRLHHLRTSQQLQPP
(54) -----DÜEAEIGHIEAEILRLSSRLHHLRTSKQSEPP
          CDS0689 Zm variantrev
            CDS0689 So variant
               CDS0689 Bn est1
                                (86) KTGKVRHGDEDAEIEEVEKENNRLSIRLESLREEKAEQIA
CDS0689 Eschscholzia californica
               CDS0689 Ga est1
                               (105) EEKKRDEKKIDMEIEEIEKEVARLSARLESLRIEKPNIMQ
                   CDS0689 Pt
                                (76) -----
      CDS0689 Plumbao zeylanica
                                (78) -----
        CDS0689 Citrus sinensis
                               (100) EEETRDERKEDIELEEIEKEISRISSRIEALREEKIDIKT
                    Consensus
                               (121)
                                            ID EI IE EI RLSSRL LRL K
```

#### FIGURE 3

## 3/10

```
401
                                     (339) IQSSVVRKRSLPENDKDESKRNDKKRSLSVGKTRVSQTES
                          CDS0689
                                     (294) TGEKDVRKRSLPEDEEKENHKRSEKR-----RASDESN (284) GGD-ARKRSFSEN----N
                       CDS0689 At
  CDS0689 Medicago trunculata
                                      (297) AAAAATAKRMAGSSKMEVIPSRYSETPGESLESSGAQERR
                      CDS0689 Os
         CDS0689 Hv contig 123
                                      (111) ISTASTCKRPAGSSKVKVPSRYSKMPGKSLG-AATQDGR
                                       (1) ------GRYSIMPGRSLGAASQERRR
(87) TSNVATTKRPAGSSKVEVVPSRYSEPPGSSLFAVTQGNRC
              CDS0689 Ta contig
             CDS0689 Zm partial
        CDS0689 Sacc sp 3'
CDS0689 Pinus taeda 3'
                                      (84) TSNAATAKRPAGSSKVRVVPSRYSETPGEYLAAVSQDKRS
                                        (1) -----
                       Consensus
                                     (401)
                                                                      RYSL PGA LG
                                            441
                                     (379) KNL---GRESRVKKRWEIPSEIVVHGNTESEKSPLSTIVK
(327) K-----SEGRVKKRWEIPSEVDLYSSGEN--GDESPIVK
                          CDS0689
                      CDS0689 At
                                     (297) KGL---GEIRAKKWEIPIEVDUSGE-----FUM--
(337) KKQSLPGSSGDANQNEEIRÄKVIËPSN--DPLSPCTSKV
(150) KESLPGSTGSTGQEEIKÄVPTEPVD--DDLSPESIDKV
(21) KESLPGSTGGAGQKEEEIKÄMPTEPVD--DDLSPESIDKV
 CDS0689 Medicago trunculata
                      CDS0689 Os
         CDS0689 Hv contig 123
             CDS0689 Ta contig
            CDS0689 Zm partial
                                     (127) KQS----LPGSATETEVNLTEPPNDE----LSPEENAKV
            CDS0689 Sacc sp 3'
                                     (124) KQS-LPGPASAASQREEIRAKLTEPSK--DELSPETVAKV
       CDS0689 Pinus taeda 3'
                                       (1) -----XEARINFGTGNSAIMAGGTKEPDTLERH
                       Consensus
                                     (441) K
                                                   GS
                                                             REÎ A E
                                                                                  LSPESL KV
                                                                Motif 4
                                            481
                                    (416) POLERERIARCVNETERISSPATEMIELEGERSFFSEDE
                         CDS0689
                                    (359) — EPKURTÜRRVGGSPEDEGAN VARLUMK-----
(324) — DEKUSTMRFVDESPRÜSGAN VARLUMK-----
(375) AEMERKURTMPPPDESPRÜSGAN VARLUGKRSFFTÄNÄ
(188) AELLERURTMFRPNETPPDSGCAN AADLVGERSFFTÄNÄ
(59) AELLERURTMPPPDETPRÜSGCAN AADLVGERSFFANÄ
(158) AELLERURTMPPPDETPRÜSGCAN VADLVGERSFFANÄ
(158) AELLERURTMPPSDESPRÜSGCAN VADLVGERSFFTÄNÄ
(161) AELLERURTMPSDESPRÜSGCAN VADLVGERSFFTÄNÄ
(162) KMGJEK KTERFTTESPRÜSGCAN VADLVGERSFFTÄNÄ
(29) KMGJEK KTERFTTESPRÜSGCAN VADLVGERSFFTÄNÄ
(29) KMGJEK KTERFTTESPRÜSGCAN VADLVGERSFFTÄNÄ
                     CDS0689 At
CDS0689 Medicago trunculata
                     CDS0689 Os
        CDS0689 Hv contig 123
             CDS0689 Ta contig
            CDS0689 Zm partial
            CDS0689 Sacc sp 3'
       CDS0689 Pinus taeda 3'
                      Consensus
                                    (481) AELLPRIRTMP DESPRDSGCAKRVADLVGKRSFF AAA
                                           521
                         CDS0689
                                    (456) -----DKEPPVCQVLSFAEEDAEEE-----
                     CDS0689 At
                                    (389) -----DRNFTFCOULKFEE-----
CDS0689 Medicago trunculata
                                    (361) EEERVMVEEEGGSVCQVLNFAEDDDDDDDDDGEQG-----
                                    CDS0689 Os
       CDS0689 Hv contig 123
            CDS0689 Ta contig
           CDS0689 Zm partial
                                    (198) DDG----NLVTPYQARVVELESPEAAAEEAEA-----
           CDS0689 Sacc ap 3'
                                    (201) BDG----NFVTPYQXPVGEL-----
      CDS0689 Pinus taeda 3'
                                     (69) ITST--PSIDXXDAGEPLERESVEEIXHAXXX-----
                      Consensus
                                    (521)
                                                            ARVLE EA E
                                           561
                                                                              592
                        CDS0689
                                    (476) -----
                    CDS0689 At
                                    (403) -----
CDS0689 Medicago trunculata
                                    (395) -----
                                    (432) -----
                    CDS0689 Os
       CDS0689 Hv contig 123
                                    (265) ALSDEAAAAAAAAEALSDEAAAAEALSDEAAA
            CDS0689 Ta contig
                                    (136) GEALGDEAAA------(226)
           CDS0689 2m partial
                                    (226)
          CDS0689 Sacc sp 3'
                                    (217)
      CDS0689 Pinus taeda 3'
                                     (99)
                                        Consensus (561)
```

## 4/10

# SEQ ID NO 1: Nicotiana tabacum seedy1 coding sequence (CDS0689)

atqaqtqttqttacaatacccaqaaqqqattqacccaqcaqatqttcaqatatggaacaatgc agcatttgataatggagattctgaagatttgtcttcgctgaaacgttcttggtctcctctga aacccctttcggttaggccatcagattcctttgaatctgatttgtcaagtaaggaaaatcaa actcctttatttgagaattcatctgttaatctctcatctccgttacccataaagccacttaa ccctaatggggctctggaaaattcaagactcaagccgaacaagcccaattccaaacagagtc ttgatgagatgqcggctagaaagaqcgqaaaqqqaaatgatttccgtgatgagaagaaaata gacgaggaaattgaagaaattcagatggagattagtaggttgagttcaagattagaggcttt qaqaattqaaaaqqctqaqaaaactqttqctaaqactqttqaaaaaqcqaqqaaqqqttqtqq caqcaaaqtttatqqaqccaaaacaaaqtqttattaaqattqaagaqcqtatatcaatgagt gcaagaacaaaggtggagcagagaggggtcttagtttaggaccatctgagatttttactgg aacgcggcggcgagggttgagtatggggccatcagatattctagcagggacaacaaaggcac ggcaattgggaaagcaagagatgattattactcctattcagccaatacaaaacaggcgaaag tcgtgtttttggaagcttcaagagattgaagaagagggaaaaagttcaagccttagtcctaa aaaqatctcqaaaaqtctqctqctaataaqaaqccccaqaqqccggggggggttgtggc tagtaggtataatcagagtacaattcagtcatcagtagtgagaaagaggtctttacctgaaa atgataaggatgagagtaagagaaatgataagaaacggtcgttatctgtagggaaaacgcgt qtqtctcaaactqaqaqcaaqaatttqqqtactqaaaqtaqqqtqaaaaaqaqatgggaaat tcctagtgagattgtagttcatggaaacacagagagtgagaaatctccactaagcattattg tgaagcctgatttgcttccgcgaattaggattgctcggtgtgtaatgagactcttagggat tctqqacctqctaaaaqaatgataqagttqataqqcaagaaatcgtttttcagtagtgatga agataaggagccacctgtctgtcaagttttaagttttgcagaggaagatgctgaagaggaa**t** aa

# SEQ ID NO 2: Nicotiana tabacum seedyl protein (CDS0689)

MSVLQYPEGIDPADVQIWNNAAFDNGDSEDLSSLKRSWSPLKPLSVRPSDSFESDLSSKENQ TPLFENSSVNLSSPLPIKPLNPNGALENSRLKPNKPNSKQSLDEMAARKSGKGNDFRDEKKI DEEIEEIQMEISRLSSRLEALRIEKAEKTVAKTVEKRGRVVAAKFMEPKQSVIKIEERISMS ARTKVEQRRGLSLGPSEIFTGTRRRGLSMGPSDILAGTTKARQLGKQEMIITPIQPIQNRRK SCFWKLQEIEEEGKSSSLSPKSRKTAARTMVTTRQAVTTIASKKNLKKDDGLLSSVQPKKLF KDLEKSAAANKKPQRPGRVVASRYNQSTIQSSVVRKRSLPENDKDESKRNDKKRSLSVGKTR VSQTESKNLGTESRVKKRWEIPSEIVVHGNTESEKSPLSIIVKPDLLPRIRIARCVNETLRD SGPAKRMIELIGKKSFFSSDEDKEPPVCQVLSFAEEDAEEE

# SEQ ID NO 3: Oryza sativa seedyl coding sequence

atggaggaggacccgctcatcccgctggtccacgtctggaacaacgccgccttcgacgactc ctcgtgttccagatcggcttggctcccccaaagccccgccgtcgcgqccqtccgcaaqqqcq acaaggagaatcaccgcccgaggttgttgatgtcgccgccggctacgacgtcgaggccgag atcggccacatcgaggcggagatcctgcgcctctcgtcccgggctccaccatctccgcgtctc caagcagccggagcccaaccgcgacgacgctccgatgggggagatggtcgcgaaggtgaggc  $\verb|cccggccgaggggcctcagcctcgggcccctggatgtgatctccatcgtcaatcgtgagaag|$ catccgctgcgcaccaagcagcctccggcgacgcggggcagggggctcagcctcgggcccat ggagatcgccgcggcgaaccctagggtgcccgcggcggcgcagcatcagcaacagcaacgcg ctggcacggcgcggatcctgaagccaatcaaggagcctccggtgcagcgtcgcaggggcgtc agectegggeegttggagateeaceaeggegteggeageaaggeaecageggeggegegage caagecgttcaccaccaagetcaacgccattcgagaagaaacccgaccetccaagcaattcg ccgtccccgccaagccatgqccqtcqaqcaatacaagqcaqacactqqactcqaqqcaagqa acagcagcaagtcgagcgaaggcgaggagcccgagccccaggcccaggagqcaatccaatgg caaggctactgacacaaggggaggcaacaaggtggtggatgagctcaagcccaaaggtgcgt cgtcaagtcagagcggcagcgccgccgccgccactgccaagaggatggcggggagctcc aagatgagggtcatcccgagccgctacagcctcactcctqqcqcttcccttqqaaqcaqtqq agcacaggagaggcgacgcaagcagtctctcccaggatcatcaggggatgcgaaccagaatg aggaaatcagagcgaaggtcatcgagccttccaatgatccactctctcctcaaacgatctcc aaggttgctgaaatgctcccaaagatcaggaccatgccgcctcctgacgagagccctcgcga  ${\tt ttccggatgcgccaagcggttgccgaattggtcgggaagcgctcgttcttcacggctgcag}$ ccgaggacgggcggctcgacgtcgaagcacccgaggcggtcgcagaagcttgagatgaa ccaccatggtttgatccgttccttccatcagctc

# SEQ ID NO 4: Oryza sativa seedyl protein

MEEDPLIPLVHVWNNAAFDDSSCSRSAWLPQSPAVAAVRKGDKENHRPEVVDVAAGYDVEAE IGHIEAEILRLSSRLHHLRVSKQPEPNRDDAPMGEMVAKVRPRPRGLSLGPLDVISIVNREK HPLRTKQPPATRGRGLSLGPMEIAAANPRVPAAAQHQQQQRAGTARILKPIKEPPVQRRRGV SLGPLEIHHGVGSKAPAAARAKPFTTKLNAIREETRPSKQFAVPAKPWPSSNTRQTLDSRQG TAASRAKARSPSPRPRRQSNGKATDTRGGNKVVDELKPKGASSSQSGSAAAAATAKRMAGSS KMRVIPSRYSLTPGASLGSSGAQERRRKQSLPGSSGDANQNEEIRAKVIEPSNDPLSPQTIS KVAEMLPKIRTMPPPDESPRDSGCAKRVAELVGKRSFFTAAAEDGRALDVEAPEAVAEA

FIGURE 4 (continued)

### 6/10

# SEQ ID NO 5: Medicago trunculata seedy1 encoding sequence

aaaaacgttaaggactaaaaatataataaaatttaagtagggattcataatggaagcacccc  ${\tt tatttacagggatcttaaatataattaaccctaatatttatgacagaaacccttttgaaatc}$ acatcggagcgtgtatgagtagccgtttcacatccaacggccagtaagagcgtaactttatt  $\verb|tcttccctcttcaatctccaacggtcacataatctcttccaaatacaaataattccctcttt|$ caacctcactcttcatttcttcaacccaaacccaaaaactaatcagattcttcttaaatct caacaacattcttcttcattccacacaggttcaagtgtggaacaacgcagcattcgatggtg aagatttcgccatgaattcatcttctgattccatcaaagagaatctaaacccatccgcattc aacattgttccttcttcaaacaaaagaactattgatgatgaaattgcggaaattgaaagtga aattaagcgattaacttcgaagctggaattgcttcqtgttgaaaaagctgaaagaaaatcg cttctgaaaagcgtgttagtggaattggtactggaagaatagtagcagcgaagtttatggaa ccgaagaaaaacgttacaccgaaacgatgtcgttttcaaggaggagacaccgaaacg aaacggtgtcgtttcggatacgccgaaatctagggttaattggagaagagggatgagtttag gtccgatggagattgccgggaaagtgatggcaccgccggcgatgacgattactccggcgacg gtgaatcggaggaagtcttgtttctggaaaccgcaggaaagttgtgaagtaatgccgtcggg gattactccggcgacggtgaataggaggaaatcttgttttttgaaacctcaagaaagttgtg aagaaaatcgaagaaaacgatttgcaaaccgaatttgaatttgaattcaaattcaqttaat  $\verb|tctgcggttggatcgattaagcgtgtgaagaagaagatgaagaaattgctcaggttcaacc|$ gaagaagctgtttgaaggtgaaaaatcagtgaagaaatcgttgaaacaaggtagaattgttg caagccggtataattccggtggtggtggtggtgatgcgaggaaaagatcgttttcggagaat aataagggtttagggagtgaaatcagggctaagaagaqatgggagataccaattgaagaagt ggatgtgagtggtttttgttatgttaccgaagatttcgacaatgaggtttgttgatgagagtc ctagagattctggtgctgttaaaagagttgctgaattgaatggaaaaagatcttacttttgt gaatttttgctgaagatgatgatgatgatgattattgtqqaacaagggtaattqtqqaaat tggaattgatttgttttgtggggttgtgtgtggaactggctatgttctgcttgattcttttgc attttggtgtgaaactaaagatgaggtgaaaagtttatgcttgttaaattggattggtttat  $\verb|tttata| at \verb|ggattta| at ttgtata| cacaatata| at at a gtatgcattgag ag ag tt$ tttcgttcagtattcattctgattttagtgtttatctcattctagaagattgtattttqttq

# SEQ ID NO 6: Medicago trunculata seedy1 protein

MNNTNNNILLHSTQVQVWNNAAFDGEDFAMNSSSDSIKENLNPSAFNIVPSSNKRTIDDEI AEIESEIKRLTSKLELLRVEKAERKIASEKRVSGIGTGRIVAAKFMEPKKNVTPKRNGVVFK EETPKRNGVVSDTPKSRVNWRRGMSLGPMEIAGKVMAPPAMTITPATVNRRKSCFWKPQESC EVMPSGITPATVNRRKSCFLKPQESCEENRRKTICKPNLNLNSNSVNSAVGSIKRVKKKDEE IAQVQPKKLFEGEKSVKKSLKQGRIVASRYNSGGGGGDARKRSFSENNKGLGSEIRAKKRWE IPIEEVDVSGFVMLPKISTMRFVDESPRDSGAVKRVAELNGKRSYFCDEDEEERVMVEEEGG SVCQVLNFAEDDDDDDDDGGOG

SEQ ID NO 8: Saccharum sp. seedyl protein (partial N-term)
MEEDPLIPLVHVWNNAAFDHASSSAWHAHSPVPASARREAEGDKENHRPDPDVEAEIGHI
EAEILRLXSRLHHLRTSKQSEPSKRGEVAPAPAAKAKAAAAARLRTRGLSLGPLDVAAAGNP
NPLTTDNQQQQPRAAQGLKPIKQATAAAGKGVRLGPLRHGRR

SEQ ID NO 9: Zea Mays seedyl encoding sequence (partial 3'end) ccacgcgtccggccgttcgagaggaggaaggccagcgttccaaggagcacgccgtccccgcc agaccgtggccatccagcaatgccaggcacccactggatgccaggcaaggcaccgcagcaag cagagccaaggcgaggagcataagccccagcaggttcaggaggcagtccacttcca gcggtcaatcacaccagcaatgtagccacgacgaagaggccggcggggagctccaaggtcag ggttgtcccgaqccgctacagcatcccacctggctcctccctagcagctgtgacacaaggca accgatgcaagcagtctctcccaggatcggctactgagaccagagtaaatctcactgagccg ccgaacgacgagttgtctcctgaagaacttgccaaggttgcagagctgctcccaaggattag gaccatgccgccttctgatgagagcccgcgtgactcgggatgtgccaagcgtgttgctgatt tggtcgggaagcgatccttcttcactgctgcaggggacgatggcaatctcgttacgccctac caggcacgggtggttgaacttgaatcacccgaggcagcagcagaagaagcagaagcttgaga agtttgtctttgatcaattccgaagtggcttgcatctgggcgtggcctctttttgcagtgtg agacattgcttagtacttttgtgttgccttgtgaaaagagagtggaaggttcatctgctgatnccttgtt

SEQ ID NO 10: Zea Mays seedyl protein (partial C-term)
TRPAVREEEGQRSKEHAVPARPWPSSNARHPLDARQGTAASRAKARSGSISPSRFRRQSTSK
AAETRAGNAKPTEATRGGSEAVNHTSNVATTKRPAGSSKVRVVPSRYSIPPGSSLAAVTQGN
RCKQSLPGSATETRVNLTEPPNDELSPEELAKVAELLPRIRTMPPSDESPRDSGCAKRVADL
VGKRSFFTAAGDDGNLVTPYOARVVELESPEAAAEEAE

SEQ ID NO 11: Arabidospis thaliana seedyl coding sequence

atgacatcaattgaggcaacagaaacgcttaacgctcctccaaagcttcagatctggaacaa cgctgccttcgacgatggagattctcaaatcacttccgccatcgaagcttcttcttggtctc  ${\tt acctcaacgaatcattcgattccgattgtagcaaggagaatcagtttccgatttcggtttcc}$ tcttcgctccaatcctcagtctcgatcaccqaaqctccgtcagcaaaatccaagaccgtgaa gaccaaatccgccgcagatcggagtaaaaaqcqaqatatcqatqcagagatcgaagaaqtag agaaggagatcggacgattatcgacgaaattgqaqtcgctccgattagagaaggcggagcaa accgcaagaagcattgctatacgtggaagaatcgttccggcgaagttcatggaatcatctca qaaacaaqtqaaattcqacqattcqtqttttacaqqatcqaaatcaaqaqccactcqtaqaq gcgttagtcttggaccagcggagatattcaattccqcqaaqaaatctgaaactqtgactcct cttcaatcagctcagaatcgacgcaagtcttgtttctttaagcttcctggaatcgaagaagg tcaagtgacgacacgaggtaaaggaagaacgagtttgagtctgagtccgagatctcgcaaag cgaaaatgacggcagctcagaagcaagcagctacgacggtggggtcaaagagagctgtgaag aaagaagaaggagttetettaacaateeageetaagaggetatteaaagaagatgaaaagaa tgtttctttaaggaaaccattgaaaccaggaagagttgtggctagtaggtacagtcaaatgg gtaaaacgcagactggagagaaagatgttaggaaaaggtcgttgcctgaggatgaagagaaa gagaatcataagaggtcggagaagagagagcttctgatgaaagtaacaagagtgaagggag agtgaagaagatgggagattccaagtgaagttgatctgtatagcagtggtgagaacggtg acgagtctcctatagttaaggagctacctaagatcagaacqcttcqtcqtqtqqqaggagc cctcgtgattcaggtgctgctaagagagttqcagaattacaagccaaggatcgtaacttcac tttttgccagcttctgaagtttgaagaatgaatgatccgcttatcaatttgagtaaaatcca caactcttqttqtqqtt

## SEQ ID NO 12: Arabidospis thaliana seedyl protein

MTSIEATETLNAPPKLQIWNNAAFDDGDSQITSAIEASSWSHLNESFDSDCSKENQFPISVS SSLQSSVSITEAPSAKSKTVKTKSAADRSKKRDIDAEIEEVEKEIGRLSTKLESLRLEKAEQ TARSIAIRGRIVPAKFMESSQKQVKFDDSCFTGSKSRATRRGVSLGPAEIFNSAKKSETVTP LQSAQNRRKSCFFKLPGIEEGQVTTRGKGRTSLSLSPRSRKAKMTAAQKQAATTVGSKRAVK KEEGVLLTIQPKRLFKEDEKNVSLRKPLKPGRVVASRYSQMGKTQTGEKDVRKRSLPEDEEK ENHKRSEKRRASDESNKSEGRVKKRWEIPSEVDLYSSGENGDESPIVKELPKIRTLRRVGGS PRDSGAAKRVAELQAKDRNFTFCQLLKFEE

# 9/10

# SEQ ID NO 13: Sequence of the [PRO0090 - CDS0689 - terminator] expression cassette

cttctacatcggcttaggtgtagcaacacgactttattattattattattattattatt attttacaaaaatataaaatagatcagtccctcaccacaagtagagcaagttggtgagttat gagtgtcaatggaacaatgaaaaccatatgacatactataattttgtttttattattgaaat tatataattcaaagagaataaatccacatagccgtaaagttctacatgtggtgcattaccaa aatatatatagettacaaaacatgacaagettagtttgaaaaattgeaateettateacatt gacacataaagtgagtgatgagtcataatattattttctttgctacccatcatgtatatatg atagccacaaagttactttgatgatgatatcaaagaacatttttaggtgcacctaacagaat atccaaataatatgactcacttagatcataatagagcatcaagtaaaactaacactctaaag caaccgatgggaaagcatctataaatagacaagcacaatgaaaatcctcatcatccttcacc acaattcaaatattatagttgaagcatagtagtaatttaaatcaactagggatatcacaagt ttgtacaaaaagcaggctggtaccggtccggaattcccgggatatcgtcgacccacgcgtc cgctgacgcgtgggttccactacatcaagacatctactactactcatcttttttgcacttatt gggtgtaaatttttgaaacccagttgagaaaaatgagtgtgttacaatacccagaagggatt gacccagcagatgttcagatatggaacaatgcagcatttgataatggagattctgaagattt gtcttcgctgaaacgttcttggtctcctctgaaacccctttcggttaggccatcagattcct ttgaatctgatttgtcaagtaaggaaaatcaaactcctttatttgagaattcatctgttaat ctctcatctccgttacccataaagccacttaaccctaatggggctctggaaaattcaagact caagccgaacaagcccaattccaaacagagtcttgatgagatggcggctagaaagagcggaa agggaaatgatttccgtgatgagaagaaatagacgaggaaattgaagaaattcagatggag attagtaggttgagttcaagattagaggctttgagaattgaaaaggctgagaaactgttgc taagactgttgaaaagcgaggaagggttgtggcagcaaagtttatggagccaaaacaaagtgcttagtttaggaccatctgagatttttactggaacgcggcggcgagggttgagtatggggc atcagatattctagcagggacaacaaaggcacggcaattgggaaagcaagagatgattatta ctcctattcagccaatacaaaacaggcgaaagtcgtgtttttggaagcttcaagagattgaa gaagagggaaaaagttcaagccttagtcctaaatcaagaaaaactgctgcaagaacaatggt tacaacaaggcaggcagttactacaattgcatcaaagaagaatttgaaaaaagatgatggac  $\verb|ttttgagttcagttcagccaaagaagtttttaaagatctcgaaaagtctgctgctaat|$ aagaagccccagaggccggggagggttgtggctagtagqtataatcagagtacaattcagtc atcagtagtgagaaagaggtctttacctgaaaatgataaggatgagagtaagagaaatgata agaaacggtcgttatctgtagggaaaacgcgtgtgtctcaaactgagagcaagaatttgggt actgaaagtagggtgaaaaagagatgggaaattcctagtgagattgtagttcatggaaacac agagagtgagaaatctccactaagcattattgtgaagcctgatttgcttccgcgaattagga ttgctcggtgtgtgaatgagactcttagggattctggacctgctaaaagaatgatagagttg  ${\tt aagttttgcagaggaagatgctgaagaggaataatgtgtaataaagggagctgctaactctt}$  ${\tt ttcatgctctttcaattttcaatcctgccttttaatttttgttcattcgtgccttttaattg}$ aatggggaagcattcttttgcttcctcaaactggtattctagcttctgaattacattgtatg agatgcggcagcggccgctctagagtatccctcgaggggcccaagcttacgcgtacccagct

## 10/10

## SEQ ID NO 14: rice prolamin RP6 promoter sequence

SEQ ID NO 15: Motif 1 CORE SEQUENCE WXNAXXD

SEQ ID NO 16: Motif 2 CORE SEQUENCE KENXXP

SEQ ID NO 17: Motif 3 (coiled coil) CORE SEQUENCE  $\text{EX}_{1-6}\text{EXXRLXXXLXXLR}$ 

SEQ ID NO 18: Motif 4 CORE SEQUENCE LPXIX<sub>1-10</sub>RDSGXXKRX<sub>1-6</sub>K

FIGURE 4 (continued)